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# UNITED STATES ENVIRONMENTAL PROTECTION AGENCY WASHINGTON, D.C. 20460

OFFICE OF PREVENTION, PESTICIDES AND TOXIC SUBSTANCES

JUN 2 5 2009

# **MEMORANDUM**

SUBJECT: BPPD Review of Monsanto's Response to SmartStax PIP DNA Analysis

(Decision No: 394799; DP Barcode: 366515).

FROM: John L. Kough, Ph.D. Senior Scientitst

Microbial Pesticides Branch, Biopesticides and

Pollution Prevention Division (7511P)

TO: Mike Mendelsohn, Senior Regulatory Action Leader

Microbial Pesticides Branch, Biopesticides and

Pollution Prevention Division (7511P)

**ACTION REQUESTED:** Review the information supplied by Monsanto to address deficiencies identified in the March 13, 2009 memorandum on DNA analysis and protein expression.

#### BACKGROUND:

The SmartStax PIP product combines several traits to control lepidopteran and coleopteran pest of corn. These traits are Cry1F (EPA reg. no. 68467-2), Cry34ab1 and Cry35Ab1 (EPA reg. no. 68467-5) from Dow AgroScience and Cry2Ab2, Cry1A.105 (EPA reg. no. 524-575) and Cry3Bb1 (EPA reg.no. 524-551) from Monsanto Company. These two combination PIP products have been traditionally bred together to provide higher levels of pest control and decrease the likelihood of developing pest insect resistance.

**CONCLUSION:** The map of plasmid PHP17661 correctly accounts for the size of the band appearing in the control lane of the Cry34Ab1 *Sac* I digest. The plasmid map and explanation of existing herbicide labels for glyphosate and glufosinate address the discrepancies identified in original submission.

CONTAINS FIFRA CONFIDENTIAL BUSINESS INFORMATION

### DATA REVIEW RECORD

Active Ingredient: Cry1a.105, Cry2Ab2, Cry1F, Cry3Bb1, Cry 34ab1, Cry35Ab1

Product Name: SmartStax corn

Company Name: Monsanto Company and Dow AgroSciences

Decision No.: 394799 DP Barcode: 366515 Submission No: 848680

## **SUMMARY OF INFORMATION SUBMITTED:**

In a letter dated April 9, 2009 from J. Austin Burns, Ph.D. representing Monsanto Company responded to the comments in a memorandum about changes in herbicide usage and discrepancies in the southern blot analysis for Cry34Ab1. The provided map of plasmid PHP17661 clarifies that the expected fragment size was 9495 bp for the *Sac* I digest for Cry34Ab1 as seen on the southern blot (lane 2, figure #7, p28 of MRID 47444904). This is more appropriate than the 13,414 bp *Sac* I fragment suggested from the plasmid map of PHP17662. The map of plasmid PHP17661 is attached.

The issue of herbicide use was addressed by the statement that there are no anticipated amendments to the glyphosate or glufosinate labels to allow a mixed use. The letter states that the only anticipated use would be a sequential use as allowed for in the existing registrations. The most likely use would be a rescue use to control weed escapes.

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